

### **In the Specification**

Please amend the paragraph on page 23 beginning at line 9 as follows:

In one embodiment, the N-terminal heterologous protein destabilization sequence is a cyclin destruction box or N-degron. In one embodiment, the C-terminal heterologous protein destabilization sequence is a CL peptide, e.g., ACKNWFSSLSHFVIHL (CL1; SEQ ID NO:89), SLISLPLPTRVKFSSLLLIRIMKIITMTFPKKLRS (CL2; SEQ ID NO:90), FYYPIWFARVLLVHYQ (CL6; SEQ ID NO:91), SNPFSSSLFGASLLIDSVSLKSNWDTSSSSCLISFFSSVMFSSTTRS (CL9; SEQ ID NO:92), CRQRFSCHLTASYPQSTVTPFLAFLRRDFFFLRHNSAD (CL10; SEQ ID NO:93), GAPHVVLFDFELRITNPLSHIQSVSLQITLIFCSLPSLILSKFLOV (CL11; SEQ ID NO:94), NTPLFKSFSTTCGVAKKTLLLAQISSLFLLSSNIAV (CL12; SEQ ID NO:95), PTVKNSPKIFCLSSSPYLA FNLEYLSLRIFSTLSKCSNTLLTSL (CL15; SEQ ID NO:96), CL216, or CL17, SNQLKRLWLWLEVRSFDRTLRRPWHLPS (CL16; SEQ ID NO:97), or SISFVIRSHASIRMGASNDFFHKLYFTKCLTSVILSKFLIHLRLRSTPRV (SL17; SEQ ID NO:98)) (see Table 1 of Gilon et al., 1998, which is specifically incorporated by reference herein), a C-ODC or a mutant C-ODC, e.g., a sequence such as HGFXXXMXXQXXGTLPMSCAQESGXXRHPAACASARINV (SEQ ID NO:81; corresponding to residues 423-461 of mODC), wherein one or more of the residues at positions marked with "X" are not the naturally occurring residue and wherein the substitution results in a decrease in the stability of a protein having that substituted sequence relative to a protein having the nonsubstituted sequence. For instance, a fusion polypeptide comprising a mutant C-ODC which has a non-conservative substitution at residues corresponding to residues 426, 427, 428, 430, 431, 433, 434, or 448 of ODC, e.g., from proline, aspartic acid or glutamic acid to alanine, can result in a fusion polypeptide with decreased stability, e.g., relative to a fusion polypeptide with a non-substituted C-ODC.